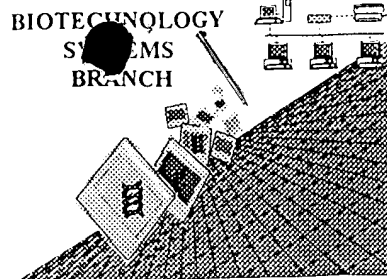


C. Myers

## RAW SEQUENCE LISTING ERROR REPORT



#13 1/2

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 08/974,584A

Source: 1655

Date Processed by STIC: 3-19-01

**THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.**

**PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:**

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

**FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.**

**FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.**

**PATENTIN 2.1 e-mail help: [patin21help@uspto.gov](mailto:patin21help@uspto.gov) or phone 703-306-4119 (R. Wax)**

**PATENTIN 3.0 e-mail help: [patin3help@uspto.gov](mailto:patin3help@uspto.gov) or phone 703-306-4119 (R. Wax)**

**TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:**

### **Checker Version 3.0**

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

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TECHNICAL CENTER  
1-800-290-2900

1655

RAW SEQUENCE LISTING DATE: 03/19/2001  
 PATENT APPLICATION: US/08/974,584A TIME: 10:11:47

Input Set : A:\-29-5us.app  
 Output Set: N:\CRF3\03192001\H974584A.raw

## SEQUENCE LISTING

4 (1) GENERAL INFORMATION:  
 6 (i) APPLICANT: Cech, Thomas R.  
 7 Lingner, Joachim  
 8 Nakamura, Toru  
 9 Chapman, Karen B.  
 10 Morin, Gregg B.  
 11 Harley, Calvin B.  
 12 Andrews, William H.  
 14 (ii) TITLE OF INVENTION: Telomerase Reverse Transcriptase  
 16 (iii) NUMBER OF SEQUENCES: 476  
 18 (iv) CORRESPONDENCE ADDRESS:  
 19 (A) ADDRESSEE: Townsend and Townsend and Crew LLP  
 20 (B) STREET: Two Embarcadero Center, Eighth Floor  
 21 (C) CITY: San Francisco  
 22 (D) STATE: California  
 23 (E) COUNTRY: USA  
 24 (F) ZIP: 94111-3834  
 26 (v) COMPUTER READABLE FORM:  
 27 (A) MEDIUM TYPE: Floppy disk  
 28 (B) COMPUTER: IBM PC compatible  
 29 (C) OPERATING SYSTEM: PC-DOS/MS-DOS  
 30 (D) SOFTWARE: PatentIn Release #1.0, Version #1.30  
 32 (vi) CURRENT APPLICATION DATA:  
 C--> 33 (A) APPLICATION NUMBER: US/08/974,584A  
 C--> 34 (B) FILING DATE: 19-Nov-1997  
 35 (C) CLASSIFICATION:  
 73 (vii) PRIOR APPLICATION DATA:  
 38 (A) APPLICATION NUMBER: US 08/724,643  
 39 (B) FILING DATE: 01-OCT-1996  
 42 (A) APPLICATION NUMBER: US 08/844,419  
 43 (B) FILING DATE: 18-APR-1997  
 46 (A) APPLICATION NUMBER: US 08/846,017  
 47 (B) FILING DATE: 25-APR-1997  
 50 (A) APPLICATION NUMBER: US 08/851,843  
 51 (B) FILING DATE: 06-MAY-1997  
 54 (A) APPLICATION NUMBER: US 08/854,050  
 55 (B) FILING DATE: 09-MAY-1997  
 58 (A) APPLICATION NUMBER: US 08/911,312  
 59 (B) FILING DATE: 14-AUG-1997  
 62 (A) APPLICATION NUMBER: US 08/912,951  
 63 (B) FILING DATE: 14-AUG-1997  
 66 (A) APPLICATION NUMBER: US 08/915,503  
 67 (B) FILING DATE: 14-AUG-1997  
 70 (A) APPLICATION NUMBER: WO PCT/US97/17618  
 71 (B) FILING DATE: 01-OCT-1997  
 74 (A) APPLICATION NUMBER: WO PCT/US97/17885

Does Not Comply  
 Corrected Diskette Needed

See p. 2

RAW SEQUENCE LISTING  
 PATENT APPLICATION: US/08/974,584A  
 DATE: 03/19/2001  
 TIME: 10:11:47

Input Set : A:\-29-Sus.app  
 Output Set: N:\CRF3\03192001\H974584A.raw

75 (B) FILING DATE: 01-OCT-1997  
 77 (viii) ATTORNEY/AGENT INFORMATION:  
 78 (A) NAME: Einhorn, Gregory P.  
 79 (B) REGISTRATION NUMBER: 38,440  
 80 (C) REFERENCE/DOCKET NUMBER: 015389-002950US  
 82 (ix) TELECOMMUNICATION INFORMATION:  
 83 (A) TELEPHONE: (415) 576-0200  
 84 (B) TELEFAX: (415) 576-0300

# ERRORED SEQUENCES

5488 (2) INFORMATION FOR SEQ ID NO: 115:  
 5490 (i) SEQUENCE CHARACTERISTICS:  
 5491 (A) LENGTH: 27 amino acids  
 5492 (B) TYPE: amino acid  
 5493 (C) STRANDEDNESS:  
 5494 (D) TOPOLOGY: linear  
 5496 (ii) MOLECULE TYPE: peptide  
 5499 (ix) FEATURE:  
 5500 (A) NAME/KEY: Modified-site  
 5501 (B) LOCATION: 18  
 5502 (D) OTHER INFORMATION: /product= "OTHER"  
 5503 /note= "Xaa = hydrophobic amino acid,  
 5504 selected from Ala, Leu, Ile, Val, Pro,  
 5505 Phe, Trp or Met"  
 5507 (ix) FEATURE:  
 5508 (A) NAME/KEY: Modified-site  
 5509 (B) LOCATION: 22  
 5510 (D) OTHER INFORMATION: /product= "OTHER"  
 5511 /note= "Xaa = hydrophobic amino acid,  
 5512 selected from Ala, Leu, Ile, Val, Pro,  
 5513 Phe, Trp or Met"

E--> 5516 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 116:  
 W--> 5518 Lys Xaa Tyr Xaa Gln Xaa Xaa Gly Ile Pro Gln Gly Ser Xaa Leu Ser  
 5519 1 5 10 15  
 W--> 5521 Xaa Xaa Leu Xaa Xaa Xaa Tyr Xaa Asp Leu  
 5522 20 25

*wrong seq i.d. number*

*115?*

## VERIFICATION SUMMARY

DATE: 03/19/2001

PATENT APPLICATION: US/08/974,584A

TIME: 10:11:51

Input Set : A:\-29-5us.app

Output Set: N:\CRF3\03192001\H974584A.raw

L:33 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]  
L:34 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]  
L:744 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:747 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:750 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:753 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:756 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:759 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
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L:771 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:774 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:777 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:780 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:783 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:786 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
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L:810 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
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L:831 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
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L:870 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
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L:879 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4

## VERIFICATION SUMMARY

DATE: 03/19/2001

PATENT APPLICATION: US/08/974,584A

TIME: 10:11:51

Input Set : A:\-29-5us.app

Output Set: N:\CRF3\03192001\H974584A.raw

L:882 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:885 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:888 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:891 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:948 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=5, Value={DNA}  
L:1598 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=22, Value={DNA}  
L:1729 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=28, Value={DNA}  
L:1745 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=29, Value={DNA}  
L:1761 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=30, Value={DNA}  
L:1777 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=31, Value={DNA}  
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L:1827 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=34, Value={DNA}  
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L:1859 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=36, Value={DNA}  
L:1875 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=37, Value={DNA}  
L:1891 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=38, Value={DNA}  
L:1907 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=39, Value={DNA}  
L:1923 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=40, Value={DNA}  
L:1939 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=41, Value={DNA}  
L:1955 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=42, Value={DNA}  
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L:1987 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=44, Value={DNA}  
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L:2036 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=47, Value={DNA}  
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L:2084 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=50, Value={DNA}  
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L:3669 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 68  
L:3681 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 68  
L:3705 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 68  
L:3709 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 68  
L:3789 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 68  
L:3805 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 68  
L:3809 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 68  
L:3841 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 68  
L:3901 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 68  
L:3905 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 68  
L:3913 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 68  
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L:4171 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=70, Value={DNA}  
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## VERIFICATION SUMMARY

PATENT APPLICATION: US/08/974,584A

DATE: 03/19/2001

TIME: 10:11:51

Input Set : A:\-29-5us.app

Output Set: N:\CRF3\03192001\H974584A.raw

L:4578 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=87, Value=[DNA]  
L:4594 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=88, Value=[DNA]  
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L:4658 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=92, Value=[DNA]  
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L:4690 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=94, Value=[DNA]  
L:4706 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=95, Value=[DNA]  
L:4722 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=96, Value=[DNA]  
L:4738 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=97, Value=[DNA]  
L:4754 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=98, Value=[DNA]  
L:4770 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=99, Value=[DNA]  
L:5087 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=102, Value=[DNA]  
L:5103 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=103, Value=[DNA]  
L:5119 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=104, Value=[DNA]  
L:5135 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=105, Value=[DNA]  
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